

## AMENDMENT

## In the claims:

Please cancel previously added claims 23-49 and replace with claims 23-49 as follows:

23. (New) A method of producing a polypeptide comprising the amino acid sequence of

Gly Pro Glu Thr Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe Val Cys Gly Asp Arg  
Gly Phe Tyr Phe Asn Lys Pro Thr Gly Tyr Gly Ser Ser Arg Arg Ala Pro Gln Thr Gly Ile  
Val Asp Glu Cys Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys Ala Pro Leu  
Lys Pro Ala Lys Ser Ala; or

Ala Tyr Arg Pro Ser Glu Thr Leu Cys Gly Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys  
Gly Asp Arg Gly Phe Tyr Phe Ser Arg Pro Ala Ser Arg Val Ser Arg Arg Ser Arg Gly Ile Val  
Glu Glu Cys Cys Phe Arg Ser Cys Asp Leu Ala Leu Leu Glu Thr Tyr Cys Ala Thr Pro Ala  
Lys Ser Glu,

comprising expressing said polypeptide in a suitable host cell transformed with a  
polynucleotide encoding said polypeptide, wherein said polynucleotide comprises a nucleic  
acid sequence selected from the group consisting of:

(a) 5'-GGA CCG GAG ACG CTC TGC GGG GCT GAG CTG GTG GAT GCT CTT  
CAG TTC GTG TGT GGA GAC AGG GGC TTT TAT TTC AAC AAG CCC ACA GGG  
TAT GGC TCC AGC AGT CGG AGG GCG CCT CAG ACA GGT ATC GTG GAT GAG  
TGC TGC TTC CGG AGC TGT GAT CTA AGG AGG CTG GAG ATG TAT TGC GCA  
CCC CTC AAG CCT GCC AAG TCA GCT-3'; and

(b) 5'-GCT TAC CGC CCC AGT GAG ACC CTG TGC GGC GGG GAG CTG  
GTG GAC ACC CTC CAG TTC GTC TGT GGG GAC CGC GGC TTC TAC TTC AGC  
AGG CCC GCA AGC CGT GTG AGC CGT CGC AGC CGT GGC ATC GTT GAG GAG  
TGC TGT TTC CGC AGC TGT GAC CTG GCC CTC CTG GAG ACG TAC TGT GCT  
ACC CCC GCC AAG TCC GAG-3'.

24. (New) A method of producing a polypeptide comprising the amino acid  
sequence of

Gly Pro Glu Thr Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe Val Cys Gly Asp Arg  
Gly Phe Tyr Phe Asn Lys Pro Thr Gly Tyr Gly Ser Ser Arg Arg Ala Pro Gln Thr Gly Ile  
Val Asp Glu Cys Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys Ala Pro Leu  
Lys Pro Ala Lys Ser Ala; or

Ala Tyr Arg Pro Ser Glu Thr Leu Cys Gly Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys  
Gly Asp Arg Gly Phe Tyr Phe Ser Arg Pro Ala Ser Arg Val Ser Arg Arg Ser Arg Gly Ile Val  
Glu Glu Cys Cys Phe Arg Ser Cys Asp Leu Ala Leu Glu Thr Tyr Cys Ala Thr Pro Ala  
Lys Ser Glu.

comprising introducing into a suitable host cell a nucleic acid molecule comprising a polynucleotide encoding said polypeptide, wherein said polynucleotide comprises a nucleic acid sequence selected from the group consisting of:

(a) 5'-GGA CCG GAG ACG CTC TGC GGG GCT GAG CTG GTG GAT GCT CTT  
CAG TTC GTG TGT GGA GAC AGG GGC TTT TAT TTC AAC AAG CCC ACA GGG  
TAT GGC TCC AGC AGT CGG AGG GCG CCT CAG ACA GGT ATC GTG GAT GAG  
TGC TGC TTC CGG AGC TGT GAT CTA AGG AGG CTG GAG ATG TAT TGC GCA  
CCC CTC AAG CCT GCC AAG TCA GCT-3'; and

(b) 5'-GCT TAC CGC CCC AGT GAG ACC CTG TGC GGC GGG GAG CTG  
GTG GAC ACC CTC CAG TTC GTC TGT GGG GAC CGC GGC TTC TAC TTC AGC  
AGG CCC GCA AGC CGT GTG AGC CGT CGC AGC CGT GGC ATC GTT GAG GAG  
TGC TGT TTC CGC AGC TGT GAC CTG GCC CTC CTG GAG ACG TAC TGT GCT  
ACC CCC GCC AAG TCC GAG-3'.

25. (New) The method of claim 23 wherein said amino acid sequence is IGF-I and  
said nucleic acid sequence is sequence (a).

26. (New) The method of claim 23 wherein said amino acid sequence is IGF-II  
and said nucleic acid sequence is sequence (b).

27. (New) A method of producing a polypeptide comprising the amino acid sequence of Fig. 1 in a suitable host cell transformed with a polynucleotide encoding said polypeptide, wherein said polynucleotide comprises the following nucleic acid sequence, wherein U can also be T:

5'-CUG GCG CUG UGC CUG CUC ACC UUC ACC AGC UCU GCC ACG GCU GGA CCG GAG ACG CUC UGC GGG GCU GAG CUG GUG GAU GCU CUU CAG UUC GUG UGU GGA GAC AGG GGC UUU UAU UUC AAC AAG CCC ACA GGG UAU GGC UCC AGC AGU CGG AGG GCG CCU CAG ACA GGU AUC GUG GAU GAG UGC UGC UUC CGG AGC UGU GAU CUA AGG AGG CUG GAG AUG UAU UGC GCA CCC CUC AAG CCU GCC AAG UCA GCU CGC UCU GUC CGU GCC CAG CGC CAC ACC GAC AUG CCC AAG ACC CAG AAG GAA GUA CAU UUG AAG AAC GCA AGU AGA GGG AGU GCA GGA AAC AAG AAC UAC AGG AUG-3',

which method comprises expressing said polynucleotide in said host cell.

28. (New) A method of producing a polypeptide comprising the amino acid sequence of Fig. 2 in a suitable host cell transformed with a polynucleotide encoding said polypeptide, wherein said polynucleotide comprises the following nucleic acid sequence, wherein U can also be T:

5'-AUG GGA AUC CCA AUG GGG AAG UCG AUG CUG GUG CUU CUC ACC UUC UUG GCC UUC GCC UCG UGC UGC AUU GCU GCU UAC CGC CCC AGU GAG ACC CUG UGC GGC GGG GAG CUG GUG GAC ACC CUC CAG UUC GUC UGU GGG GAC CGC GGC UUC UAC UUC AGC AGG CCC GCA AGC CGU GUG AGC CGU CGC AGC CGU GGC AUC GUU GAG GAG UGC UGU UUC CGC AGC UGU GAC CUG GCC CUC CUG GAG ACG UAC UGU GCU ACC CCC GCC AAG UCC GAG AGG GAC GUG UCG ACC CCU CCG ACC GUG CUU CCG GAC AAC UUC CCC AGA UAC CCC GUG GGC AAG UUC UUC CAA UAU GAC ACC UGG AAG CAG UCC ACC CAG CGC CUG CGC AGG GGC CUG CCU GCC CUC CUG CGU GCC CGC CGG GGU CAC GUG CUC GCC AAG GAG CUC GAG GCG UUC AGG GAG GCC AAA CGU CAC CGU CCC CUG AUU GCU CUA CCC ACC CAA GAC CCC GCC CAC GGG GGC GCC CCC CCA GAG AUG GCC AGC AAU CGG AAG UGA-3',

which method comprises expressing said polynucleotide in said host cell.

29. (New) The method of claim 24 wherein said nucleic acid molecule is the plasmid phigf1.

30. (New) The method of claim 24 wherein said nucleic acid molecule is the plasmid phigf2.

31. (New) A method of producing a polypeptide comprising the amino acid sequence of:

Gly Pro Glu Thr Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe Val Cys Gly Asp Arg  
Gly Phe Tyr Phe Asn Lys Pro Thr Gly Tyr Gly Ser Ser Arg Arg Ala Pro Gln Thr Gly Ile  
Val Asp Glu Cys Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys Ala Pro Leu  
Lys Pro Ala Lys Ser Ala; or

Ala Tyr Arg Pro Ser Glu Thr Leu Cys Gly Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys  
Gly Asp Arg Gly Phe Tyr Phe Ser Arg Pro Ala Ser Arg Val Ser Arg Arg Ser Arg Gly Ile Val  
Glu Glu Cys Cys Phe Arg Ser Cys Asp Leu Ala Leu Leu Glu Thr Tyr Cys Ala Thr Pro Ala  
Lys Ser Glu,

comprising expressing the heterologous DNA in the transformed cellular hosts of a composition of claim 10 or 11.

32. (New) The method of producing a polypeptide according to claim 31 wherein said amino acid sequence is IGF-I and said nucleic acid sequence is sequence (a).

33. (New) A method of producing a polypeptide according to claim 31 wherein said amino acid sequence is IGF-II and said nucleic acid sequence is sequence (b).

34. (New) A method of producing a polypeptide according to claim 31 wherein said amino acid sequence is the amino acid sequence of Fig. 1 and said composition comprises the nucleic acid sequence:

5'-CTG GCG CTG TGC CTG CTC ACC TTC ACC AGC TCT GCC ACG GCT  
GGA CCG GAG ACG CTC TGC GGG GCT GAG CTG GTG GAT GCT CTT CAG TTC  
GTG TGT GGA GAC AGG GGC TTT TAT TTC AAC AAG CCC ACA GGG TAT GGC  
TCC AGC AGT CGG AGG GCG CCT CAG ACA GGT ATC GTG GAT GAG TGC TGC  
TTC CGG AGC TGT GAT CTA AGG AGG CTG GAG ATG TAT TGC GCA CCC CTC  
AAG CCT GCC AAG TCA GCT CGC TCT GTC CGT GCC CAG CGC CAC ACC GAC  
ATG CCC AAG ACC CAG AAG GAA GTA CAT TTG AAG AAC GCA AGT AGA GGG  
AGT GCA GGA AAC AAG AAC TAC AGG ATG-3'.

35. (New) A method of producing a polypeptide according to claim 31, wherein said amino acid sequence is the amino acid sequence of Fig. 2 and said composition comprises the nucleic acid sequence:

5'-ATG GGA ATC CCA ATG GGG AAG TCG ATG CTG GTG CTT CTC ACC  
TTC TTG GCC TTC GCC TCG TGC ATT GCT GCT TAC CGC CCC AGT GAG  
ACC CTG TGC GGC GGG GAG CTG GTG GAC ACC CTC CAG TTC GTC TGT GGG  
GAC CGC GGC TTC TAC TTC AGC AGG CCC GCA AGC CGT GTG AGC CGT CGC  
AGC CGT GGC ATC GTT GAG GAG TGC TGT TTC CGC AGC TGT GAC CTG GCC  
CTC CTG GAG ACG TAC TGT GCT ACC CCC GCC AAG TCC GAG AGG GAC GTG  
TCG ACC CCT CCG ACC GTG CTT CCG GAC AAC TTC CCC AGA TAC CCC GTG  
GGC AAG TTC TTC CAA TAT GAC ACC TGG AAG CAG TCC ACC CAG CGC CTG  
CGC AGG GGC CTG CCT GCC CTC CTG CGT GCC CGC CGG GGT CAC GTG CTC  
GCC AAG GAG CTC GAG GCG TTC AGG GAG GCC AAA CGT CAC CGT CCC CTG  
ATT GCT CTA CCC ACC CAA GAC CCC GCC CAC GGG GGC GCC CCC CCA GAG  
ATG GCC AGC AAT CGG AAG TGA-3'.

36. (New) The method of claim 31 wherein said heterologous DNA molecule is located on a plasmid that replicates in said host cells.

37. (New) The method of claim 31 wherein said transformed cellular hosts are yeast.

38. (New) The method of claim 31 wherein said transformed cellular hosts are *E. coli*.

39. (New) The method of claim 31 wherein said transformed cellular hosts are *B. subtilis*.

40. (New) The method of claim 31 wherein said transformed cellular hosts are *E. coli* strain HB 101 (phigf1).

41. (New) The method of claim 31 wherein said transformed cellular hosts are *E. coli* strain HB 101 (phigf2).

42. (New) A method according to claim 23 wherein the polypeptide is IGF-I and the polynucleotide sequence is sequence (a).

43. (New) A method according to claim 23 wherein the polypeptide is IGF-II and the polynucleotide sequence is sequence (b).

44. (New) A vector comprising a nucleic acid sequence selected from the group consisting of the following nucleic acid sequences:

(a) 5'-GGA CCG GAG ACG CUC UGC GGG GCU GAG CUG GUG GAU GCU CUU CAG UUC GUG UGU GGA GAC AGG GGC UUU UAU UUC AAC AAG CCC ACA GGG UAU GGC UCC AGC AGU CGG AGG GCG CCU CAG ACA GGU AUC GUG GAU GAG UGC UGC UUC CGG AGC UGU GAU CUA AGG AGG CUG GAG

AUG UAU UGC GCA CCC CUC AAG CCU GCC AAG UCA GCU-3', wherein U can also be T;

(b) 5'-GCU UAC CGC CCC AGU GAG ACC CUG UGC GGC GGG GAG CUG GUG GAC ACC CUC CAG UUC GUC UGU GGG GAC CGC GGC UUC UAC UUC AGC AGG CCC GCA AGC CGU GUG AGC CGU CGC AGC CGU GGC AUC GUU GAG GAG UGC UGU UUC CGC AGC UGU GAC CUG GCC CUC CUG GAG ACG UAC UGU GCU ACC CCC GCC AAG UCC GAG-3', wherein U can also be T;

(c) a nucleic acid sequence complementary to (a) or (b);

(d) a fragment of (a) or (b) that is at least 18 bases in length; and

(e) a fragment of (c) that is at least 18 bases in length.

45. (New) A vector according to claim 44 wherein said nucleic acid sequence is nucleic acid sequence (a).

46. (New) An expression vector comprising a polynucleotide encoding a polypeptide, wherein said polypeptide comprises the amino acid sequence of:

(a) Gly Pro Glu Thr Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe Val Cys Gly Asp Arg Gly Phe Tyr Phe Asn Lys Pro Thr Gly Tyr Gly Ser Ser Ser Arg Arg Ala Pro Gln Thr Gly Ile Val Asp Glu Cys Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys Ala Pro Leu Lys Pro Ala Lys Ser Ala;

(b) Ala Tyr Arg Pro Ser Glu Thr Leu Cys Gly Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys Gly Asp Arg Gly Phe Tyr Phe Ser Arg Pro Ala Ser Arg Val Ser Arg Arg Ser Arg Gly Ile Val Glu Glu Cys Cys Phe Arg Ser Cys Asp Leu Ala Leu Glu Thr Tyr Cys Ala Thr Pro Ala Lys Ser Glu; or

(c) a fragment of said amino acid sequence (a) or (b),

wherein said amino acid sequence (a) is encoded by the nucleic acid sequence

5'-GGA CCG GAG ACG CUC UGC GGG GCU GAG CUG GUG GAU GCU CUU CAG UUC GUG UGU GGA GAC AGG GGC UUU UAU UUC AAC AAG CCC ACA GGG UAU GGC UCC AGC AGU CGG AGG GCG CCU CAG ACA GGU AUC GUG

GAU GAG UGC UGC UUC CGG AGC UGU GAU CUA AGG AGG CUG GAG AUG

UAU UGC GCA CCC CUC AAG CCU GCC AAG UCA GCU-3', wherein U can also be T;

wherein said amino acid sequence (b) is encoded by the nucleic acid sequence

5'-GCU UAC CGC CCC AGU GAG ACC CUG UGC GGC GGG GAG CUG GUG GAC

ACC CUC CAG UUC GUC UGU GGG GAC CGC GGC UUC UAC UUC AGC AGG CCC

GCA AGC CGU GUG AGC CGU CGC AGC CGU GGC AUC GUU GAG GAG UGC

UGU UUC CGC AGC UGU GAC CUG GCC CUC CUG GAG ACG UAC UGU GCU ACC

CCC GCC AAG UCC GAG-3', wherein U can also be T;

and wherein said fragment is encoded by a nucleic acid sequence of at least eighteen bases of  
a nucleic acid sequence selected from the group consisting of:

5'-GCU UAC CGC CCC AGU GAG ACC CUG UGC GGC GGG GAG CUG GUG GAC

ACC CUC CAG UUC GUC UGU GGG GAC CGC GGC UUC UAC UUC AGC AGG CCC

GCA AGC CGU GUG AGC CGU CGC AGC CGU GGC AUC GUU GAG GAG UGC

UGU UUC CGC AGC UGU GAC CUG GCC CUC CUG GAG ACG UAC UGU GCU ACC

CCC GCC AAG UCC GAG-3', wherein U can also be T; and

5'-GGA CCG GAG ACG CUC UGC GGG GCU GAG CUG GUG GAU GCU CUU CAG

UUC GUG UGU GGA GAC AGG GGC UUU UAU UUC AAC AAG CCC ACA GGG

UAU GGC UCC AGC AGU CGG AGG GCG CCU CAG ACA GGU AUC GUG GAU

GAG UGC UGC UUC CGG AGC UGU GAU CUA AGG AGG CUG GAG AUG UAU

UGC GCA CCC CUC AAG CCU GCC AAG UCA GCU-3', wherein U can also be T.

47. (New) An expression vector according to claim 46, wherein said amino acid sequence is the sequence of Fig. 1.

48. (New) An expression vector according to claim 46, wherein said amino acid sequence is the sequence of Fig.2.

49. (New) A composition according to claim 1 wherein said nucleic acid molecules comprise a nucleic acid sequence selected from the group consisting of (a), (b), and (d).